OM nucleic - nucleic search, using sw model

Run on: December 12, 2005, 02:19:10 ; Search time 10232 Seconds
                                              (without alignments)
                                   11110.917 Million cell updates/sec

Title: US-10-782-002-34
Perfect score: 2000
Sequence: 1 cccggtctggagacggggag..........gcctcaagttttttttcgttc 2000

Scoring table: IDENTITY_NUC
                Gapop 10.0 , Gapext 1.0

Searched: 5883141 seqs, 28421725653 residues

Total number of hits satisfying chosen parameters: 11766282

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
                 Maximum Match 100%
                 Listing first 45 summaries

Database :   GenEmbl:*
                1: gb_ba:*
                2: gb_in:*
                3: gb_env:*
                4: gb_om:*
                5: gb_ov:*
                6: gb_pat:*
                7: gb_ph:*
                8: gb_pr:*
                9: gb_ro:*
                10: gb_sts:*
                11: gb_sy:*
                12: gb_un:*
                13: gb_vi:*
                14: gb_htg:*
                15: gb_pl:*

OM nucleic - nucleic search, using sw model

Run on: December 12, 2005, 02:10:15 ; Search time 1221 Seconds
                                              (without alignments)
                                   10916.777 Million cell updates/sec

Title: US-10-782-002-34
Perfect score: 2000
Sequence: 1 cccggtctggagacggggag..........gcctcaagttttttttcgttc 2000

Scoring table: IDENTITY_NUC
                Gapop 10.0 , Gapext 1.0

Searched: 4996997 seqs, 3332346308 residues

Total number of hits satisfying chosen parameters: 9993994

```
Minimum.DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
                 Maximum Match 100%
                 Listing first 45 summaries

Database :        N_Geneseq_21:*
                  1:  geneseqn1980s:*
                  2:  geneseqn1990s:*
                  3:  geneseqn2000s:*
                  4:  geneseqn2001as:*
                  5:  geneseqn2001bs:*
                  6:  geneseqn2002as:*
                  7:  geneseqn2002bs:*
                  8:  geneseqn2003as:*
                  9:  geneseqn2003bs:*
                  10:  geneseqn2003cs:*
                  11:  geneseqn2003ds:*
                  12:  geneseqn2004as:*
                  13:  geneseqn2004bs:*
                  14:  geneseqn2005s:*


                        GenCore version 5.1.6
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OM nucleic - nucleic search, using sw model

Run on:           December 12, 2005, 02:11:55 ; Search time 8166 Seconds
                                         (without alignments)
                                         11458.996 Million cell updates/sec

Title:            US-10-782-002-34
Perfect score:    2000
Sequence:         1 cccggtctggagacgggggag..........gcctcaagttttttttcgttc 2000

Scoring table:    IDENTITY_NUC
                  Gapop 10.0 , Gapext 1.0

Searched:         41078325 seqs, 23393541228 residues

Total number of hits satisfying chosen parameters:        82156650

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
                 Maximum Match 100%
                 Listing first 45 summaries

Database :        EST:*
                  1:  gb_est1:*
                  2:  gb_est2:*
                  3:  gb_est3:*
                  4:  gb_htc:*
                  5:  gb_est4:*
                  6:  gb_est5:*
                  7:  gb_est6:*
                  8:  gb_est7:*
                  9:  gb_gss1:*
                  10:  gb_gss2:*
                  11:  gb_gss3:*


                        GenCore version 5.1.6
                Copyright (c) 1993 - 2005  Compugen Ltd.
```